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**Fig. 1****SEQ ID NO: 1**

GCCAGCCGCAGACCGTGGCTGAGCATGGAGCTGTCCCCCGCAGTCCT  
CCGGAGATGCTGGAGGAGTCGGATTGCCCGTCACCCCTGGAGCTGAAG  
TCAGCCCCCAGCAAGAAGATGTGGATTAAGCTTCGGTCTCTGCTGCGC  
TACATGGTGAAGCAGTTGGAGAATGGGGAGATAAACATTGAGGAGCTG  
AAGAAAAATCTGGAGTACACAGCTTCTCTGCTGGAAGCCGTCTACATA  
GATGAGACACGGCAAATCTTGGACACGGAGGACGAGCTGCAGGAGCTG  
CGGTCAGATGCCGTGCCTTCGGAGGTGCGGGACTGGCTGGCCTCCACC  
TTCACCCAGCAGGCCCGGGCCAAAGGCCGCCGAGCAGAGGAGAAGCCC  
AAGTTCCGAAGCATTGTGCACGCTGTGCAGGCTGGGATCTTCGTGGAA  
CGGATGTTCCGGAGAACATACACCTCTGTGGGCCCCACTTACTCTACT  
GCGGTTCTCAACTGTCTCAAGAACCTGGATCTCTGGTGCTTTGATGTC  
TTTTCTTTGAACCAGGCAGCAGATGACCATGCCCTGAGGACCATTGTT  
TTTGAGTTGCTGACTCGGCATAACCTCATCAGCCGCTTCAAGATTCCC  
ACTGTGTTTTTGTATGAGTTTCTGGATGCCTTGGAGACAGGCTATGGG  
AAGTACAAGAATCCTTACCACAACCAGATCCACGCAGCCGATGTTACC  
CAGACAGTCCATTGCTTCTTGCTCCGCACAGGGATGGTGCCTGCTG  
TCGGAGATTGAGCTCCTGGCCATCATCTTTGCTGCAGCTATCCATGAT  
TATGAGCACACGGGCACTACCAACAGCTTCCACATCCAGACCAAGTCA  
GAATGTGCCATCGTGTACAATGATCGTTCAGTGCTGGAGAATCACCAC  
ATCAGCTCTGTTTTCCGATTGATGCAGGATGATGAGATGAACATTTTC  
ATCAACCTCACCAAGGATGAGTTTGTAGAACTCCGAGCCCTGGTCATT  
GAGATGGTGTGGCCACAGACATGTCTGCCATTTCCAGCAAGTGAAG  
ACCATGAAGACAGCCTTGCAACAGCTGGAGAGGATTGACAAGCCCAAG  
GCCCTGTCTCTACTGCTCCATGCTGCTGACATCAGCCACCCAACCAAG  
CAGTGGTTGGTCCACAGCCGTTGGACCAAGGCCCTCATGGAGGAATTC  
TTCCGTCAGGGTGACAAGGAGGCAGAGTTGGGGCCTGCCCTTTTCTCCA  
CTCTGTGACCGCACTTCCACTCTAGTGGCACAGTCTCAGATAGGGTTC  
ATCGACTTCATTGTGGAGCCACATTCTCTGTGCTGACTGACGTGGCA  
GAGAAGAGTGTTTCAGCCCCCTGGCGGATGAGGACTCCAAGTCTAAAAAC  
CAGCCCAGCTTTCAGTGGCGCCAGCCCTCTCTGGATGTGGAAGTGGGA  
GACCCCAACCCTGATGTGGTTCAGCTTTCGTTCCACCTGGGTCAAGCGC  
ATTTCAGGAGAACAAGCAGAAATGGAAGGAACGGGCAGCAAGTGGCATC  
ACCAACCAGATGTCCATTGACGAGCTGTCCCCCTGTGAAGAAGAGGCC  
CCCCCATCCCCCTGCCGAAGATGAACACAACCAGAATGGGAATCTGGAT  
TAGCCCTGGGGCTGGCCCAGGTCTTCATTGAGTCCAAAGTGTTTGATG  
TCATCAGCACCATCCATCAGGACTGGCTCCCCCATCTGCTCCAAGGGA  
GCGTGGTTCGTGGAAGAAACAACCCACCTGAAGGCCAAATGCCAGAGAT  
TTGGGGTTGGGGAAAGGGCCCCCTCCCCACCTGACACCCACTGGGGTGC  
ACTTTAATGTTCCGGCAGCAAGACTGGGGAACTTCAGGCTCCCAGTGG  
TCACTGTGCCCATCCCTCAGCCTCTGGATTCTCTTCATGGCCAGGTGG  
CTGCCAGGGAGCGGGGAGCTTCTTGGAGGCTTCCCAGGGCCTTGGGGA

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**Fig. 1 (continued)**

AGGGTCAGAGATGCCAGCCCCCTGGGACCTCCCCCATCCTTTTTGCCT  
CCAAGTTTCTAAGCAATACATTTTGGGGGTTCCTCAGCCCCCCCACCC  
CAGATCTTAGCTGGCAGGTCTGGGTGCCCTTTTCCTCCCCTGGGAAG  
GGCTGGAATAGGATAGAAAGCTGGGGGTTTTCAGAGCCCTATGTGTGG  
GGAGGGGAGTGGATTCCTTCAGGGCATGGTACCTTTCTAGGATCTGGG  
AATGGGGTGGAGAGGACATCCTCTTCACCCAGAATTGCGGGAATTC

**Fig. 2**SEQ ID NO: 2

MELSPRSPPEMLEESDCPSPLELKSAPSKKMWIKLRSLRLRYMVKQLEN  
GEINIEELKKNLEYTASLLEAVYIDETRQILDTEDELQELRSDAVPSE  
VRDWLASTFTQARAKGRRAEEKPKFRSIVHAVQAGIFVERMFRRTYT  
SVGPTYSTAVLNCLKNLDLWCFDVFSLNQAADDHALRTIVFELLTRHN  
LISRFKIPTVFLMSFLDALETGYGKYKNPYHNQIHAADVTQTVHCFL  
RTGMVHCLSEIELLAIIFAAAIHDYEHTGTTNSFHIQTKSECAIVYND  
RSVLENHHISSVFRLMQDDEMNI FINLTKDEFVELRALVIEMVLATDM  
SCHFQQVKTMTALQQLERIDKPKALSLLLHAADISHPTKQWLVS  
TKALMEEFFRQGDKEAELGLPFSPLCDRTSTLVAQSQIGFIDFIVEPT  
FSVLTDVAEKSVQPLADEDSKSKNQPSFQWRQPSLDVEVGDPNPDVVS  
FRSTWVKRIQENKQKWKERAASGITNQMSIDELSPCEEEAPPSPAEDE  
HNQNGNLD

**Fig. 3**SEQ ID NO: 3

5'- TGTGGAAGTGGGAGACCCC -3'

**Fig. 4**SEQ ID NO: 4

5'- CTGAATGCGCTTGACCCAG -3'

**Fig. 5**SEQ ID NO: 5

5'- ACCCTGATGTGGTCAGCTTTCGTTCCA -3'